

AMENDMENTS TO THE SPECIFICATION

Please enter the following amendments to the Specification.

[At page 8, line 17]:

In general, variants will be at least 75% homologous, more suitably at least 80%, preferably at least 85%, and more preferably at least 90% homologous to a polypeptide as for example shown in SEQ ID NO: 2, 4 or 8.

Variant peptides or polypeptides, resulting from rational or established methods of mutagenesis or from combinatorial chemistries as hereinafter described, may comprise conservative amino acid substitutions. Exemplary conservative substitutions in an immuno-interactive polypeptide or polypeptide fragment according to the invention may be made according to the following table:

TABLE A

Original Residue	Exemplary Substitutions
Ala	Ser
Arg	Lys
Asn	Gln, His
Asp	Glu
Cys	Ser
Gln	Asn
Glu	Asp
Gly	Pro
His	Asn, Gln
Ile	Leu, Val
Leu	Ile, Val
Lys	Arg, Gln, Glu
Met	Leu, Ile,
Phe	Met, Leu, Tyr
Ser	Thr
Thr	Ser
Trp	Tyr
Tyr	Trp, Phe
Val	Ile, Leu

[Following page 28]:

Please Insert the Sequence Listing (pages 1-9)